























```

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                  /usedbio-X96381-ERR_rRNA
                  /function="transcription factor"
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ORIGIN
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Ratio: 1.000 Gaps: 0
Percent Similarity: 100.000 Percent Identity: 100.000
Alignment_Block:
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Align seg 1/1 to: HSEBM12 from: 1 to: 163

294 LYSASATGPGCAIAGCAGTAACTAAGTAAGTAAAGTGAATGGTGTTCACGCGCCCTTT
|||||
57 AAGAACGCCGCACCACGACGACTGATGACACGCTGACGCGCCCTCTT 98
seq_name: gb_pat2:E06024
seq_documentation_block:
LOCUS E06024 473 bp RNA PAT 29-SEP-1997
DEFINITION cdna encoding a protein(EIA-F) which binds to enhancer core region
ACCESSION E06024
VERSIONS E06024.1 GI:2174211
KEYWORDS JP 1993128975-A/L.
SOURCE Homo sapiens.
ORGANISM Homo sapiens.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Homidae; Homo.
REFERENCE 1 (bases 1 to 473)
AUTHORS Isono,F., Yoshida,K. and Fujinaga,M.
JOURNAL PNAS USA 90(1993)12897-12901
PATENT: JP 1993128975-A 1 14-DEC-1993;
TAKARA SHUZO CO LTD
COMMENT OS Homo sapiens (human)
PN JP 1993128975-A/1
PD 14-DEC-1993
PF 02-JUN-1993 JP 1992165453
PT F TONO FUJINAGA, KOSHIDA KOICHI, FUJINAGA MEGUMI PC C12N15/34;
CC topology: linear;
CC hypochetical: No;
CC anti-sense: No;
CC *source: cell_line=HeLa cells;
FH Key Location/Qualifiers
FT FT COS 1..468
FT FT enhancer core
FT FT region of adenovirus EIA gene' FT 3'UTR
FEATURES
source Location/Qualifiers
BASE COUNT 91 a 143 c 140 g 99 t
ORIGIN
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Quality: 14.00 Length: 14
Ratio: 1.000 Gaps: 0
Percent Similarity: 100.000 Percent Identity: 100.000
Alignment_Block:
US-09-126-945B-2.COPY.2..335 x E06024 ..

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Align seg 1/1 to: E06024 from: 1 to: 473

294 lysAsnArgProAlaMetAsnTyrAspLysLeuSerArgSer 307

166 AAGAACCGGCCAGCCATGAATTACGACCAAGCTGAGCGCTCG 207